

Application No. 09/919,162  
Amendment dated November 15, 2005  
After Final Office Action of August 23, 2005

Docket No.: LUD 5684.2 CIP (10106926)

**AMENDMENTS TO THE SPECIFICATION**

Page 6, lines 10-13, please amend as follows:

The amino acid sequence of the extracellular domain of human IL-10R was used to screen the database of the Sanger Center (~~http://www.sanger.ac.uk/cgi-bin/nph-Blast\_server.html~~), ~~incorporated by reference~~, using TBLASTN software.

Page 6, lines 18-21: please amend as follows:

Once the BAC sequence was identified, it was analyzed further, using the NIX analysis program ~~found at~~ ~~http://www.hemp.mrc.ac.uk/Registered/Webapp/nix/~~, ~~incorporated by reference~~. The software predicted a gene comprising 5 exons, stretching over about 16 kilobases, with the last exon corresponding to several EST sequences.